

EAST Search History

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L31	1192	methods of mutation identification, DNA, hybridization, probe, gene chip	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:54
L32	472	L31 and @ay<="2000"	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:54
L33	337	L31 and @ay<="1999"	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:54
L34	50	L31 and @ay<="1996"	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:54
L35	34	L31 and @ay<="1995"	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:54
L36	0	L33 and computer and hybridization and intensity	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:55
L37	860	L31 and computer and hybridization	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:55
L38	501	L37 and intensity	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:55
L39	445	L38 and array and probe	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:55
L40	140	L39 and @ay<="2000"	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:56
L41	95	L39 and @ay<="1999"	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:56
L42	53	L39 and @ay<="1998"	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:56
L43	17	L39 and @ay<="1997"	US-PGPUB; USPAT; DERWENT	AND	ON	2006/03/08 18:56

3/8/06

```
125 S10
10839557 RATIO
S11 114 S10 AND RATIO
? s11 and wild(n4)type
Processing
Processed 10 of 50 files ...
Processing
Processed 20 of 50 files ...
Processing
Processed 30 of 50 files ...
Processing
Processed 40 of 50 files ...
Processing
Processing
Processing
Processing
Processed 50 of 50 files ...
Completed processing all files
39935393 11
3326017 WILD
40805971 TYPE
1118780 WILD(4N)TYPE
S12 177612 11 AND WILD(N4)TYPE
? ds
```

Set	Items	Description
S1	20335	MUTATION(S)IDENTIF? AND SEQUENCE AND PROBE AND PATTERN
S2	0	S1 AND MISMATCH AND WILD-TYPE
S3	7318	S1 AND MISMATCH
S4	7100	S3 AND HYBRIDIZATION
S5	6058	S4 AND AFFINITY
S6	4528	S5 AND DATABASE
S7	156	S6 AND IDENTIFY AND PROBE(N5) INTENSITY
S8	145	S7 AND POSITION
S9	144	S8 AND REFERENCE
S10	125	S9 AND REFERENCE(S) SEQUENCE
S11	114	S10 AND RATIO
S12	177612	11 AND WILD(N4)TYPE

```
? s s11 and DNA(s)sequence
Processing
Processed 20 of 50 files ...
Processing
Processed 30 of 50 files ...
Processing
Processed 40 of 50 files ...
Completed processing all files
```

```
114 S11
7246795 DNA
6842594 SEQUENCE
1278932 DNA(S)SEQUENCE
S13 113 S11 AND DNA(S)SEQUENCE
```

```
?
PLEASE ENTER A COMMAND OR BE LOGGED OFF IN 5 MINUTES
? ds
```

Set	Items	Description
S1	20335	MUTATION(S)IDENTIF? AND SEQUENCE AND PROBE AND PATTERN
S2	0	S1 AND MISMATCH AND WILD-TYPE
S3	7318	S1 AND MISMATCH
S4	7100	S3 AND HYBRIDIZATION
S5	6058	S4 AND AFFINITY

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S6      4528   S5 AND DATABASE
S7      156    S6 AND IDENTIFY AND PROBE(N5) INTENSITY
S8      145    S7 AND POSITION
S9      144    S8 AND REFERENCE
S10     125    S9 AND REFERENCE(S) SEQUENCE
S11     114    S10 AND RATIO
S12     177612 11 AND WILD(N4) TYPE
S13     113    S11 AND DNA(S) SEQUENCE
? s s13 base(n5)position
>>>Term "BASE" in invalid position
? s s13 and base(n5)position
Processing
Processed 20 of 50 files ...
Processing
Processed 30 of 50 files ...
Processing
Processing
Processed 40 of 50 files ...
Processing
Completed processing all files
      113   S13
    12882015 BASE
    16049460 POSITION
      184372 BASE(5N) POSITION
S14      67   S13 AND BASE(N5) POSITION
? s s14 and point(n5)mutation
Processed 30 of 50 files ...
Processing
Processing
Processed 40 of 50 files ...
Completed processing all files
      67   S14
    20353330 POINT
    2041392  MUTATION
      168368 POINT(5N) MUTATION
S15      21   S14 AND POINT(N5) MUTATION

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